

1/31/06

```

1835383  PEPTIDE
        10048224  PROTEIN
        448889   PEPTIDE(S) PROTEIN
        4789130  BINDING
        1376400  AFFINITY
        4396852  SEQUENCE
        S1      17008  PEPTIDE(S) PROTEIN AND BINDING AND AFFINITY AND SEQUENCE
? s MHC(w)binding
        255476   MHC
        4789130  BINDING
        S2      2487   MHC(W) BINDING
? s s1 and s2
        17008    S1
        2487     S2
        S3      122    S1 AND S2
? s s3 and scal?
Sending Break...
?s s3 and profile and scoring
        122      S3
        3281498  PROFILE
        901941   SCORING
        S4      0      S3 AND PROFILE AND SCORING
? s profile and scoring
        3281498  PROFILE
        901941   SCORING
        S5      25240  PROFILE AND SCORING
? s s2 and s5
        2487     S2
        25240     S5
        S6      9      S2 AND S5
? s s3 and scoring
        122      S3
        901941   SCORING
        S7      4      S3 AND SCORING
? s s7 and s6
        4        S7
        9        S6
        S8      0      S7 AND S6
? s s7 or s6
        4        S7
        9        S6
        S9      13     S7 OR S6
? rd

```

>>>Duplicate detection is not supported for File 229.

>>>Duplicate detection is not supported for File 280.

>>>Records from unsupported files will be retained in the RD set.

S10 5 RD (unique items)

? t s10/ti/all

>>>No matching display code(s) found in file(s): 229

10/TI/1 (Item 1 from file: 5)

DIALOG(R)File 5:(c) 2006 BIOSIS. All rts. reserv.

Prediction of MHC class I binding peptides using profile motifs

10/TI/2 (Item 1 from file: 34)

DIALOG(R)File 34:(c) 2006 Inst for Sci Info. All rts. reserv.

Title: Enhancement to the RANKPEP resource for the prediction of peptide binding to MHC molecules using profiles

10/TI/3 (Item 1 from file: 73)

DIALOG(R)File 73:(c) 2006 Elsevier Science B.V. All rts. reserv.

Use of global amino acid replacements to define the requirements for MHC binding and T cell recognition of moth cytochrome c (93-103)

10/TI/4 (Item 1 from file: 98)

DIALOG(R)File 98:(c) 2005 The HW Wilson Co. All rts. reserv.

Rapid determination of HLA B*07 ligands from the West Nile virus NY99 genome.

10/TI/5 (Item 1 from file: 149)

DIALOG(R)File 149:(c) 2006 The Gale Group. All rts. reserv.

Identification of naturally processed HLA-A2-restricted proinsulin epitopes by reverse immunology. (IMMUNOLOGY AND TRANSPLANTATION)

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